

662707" FILED AT 60

PEPTIDE STRUCTURE of: human\_prot.pep ck: 2321, 1 to 458

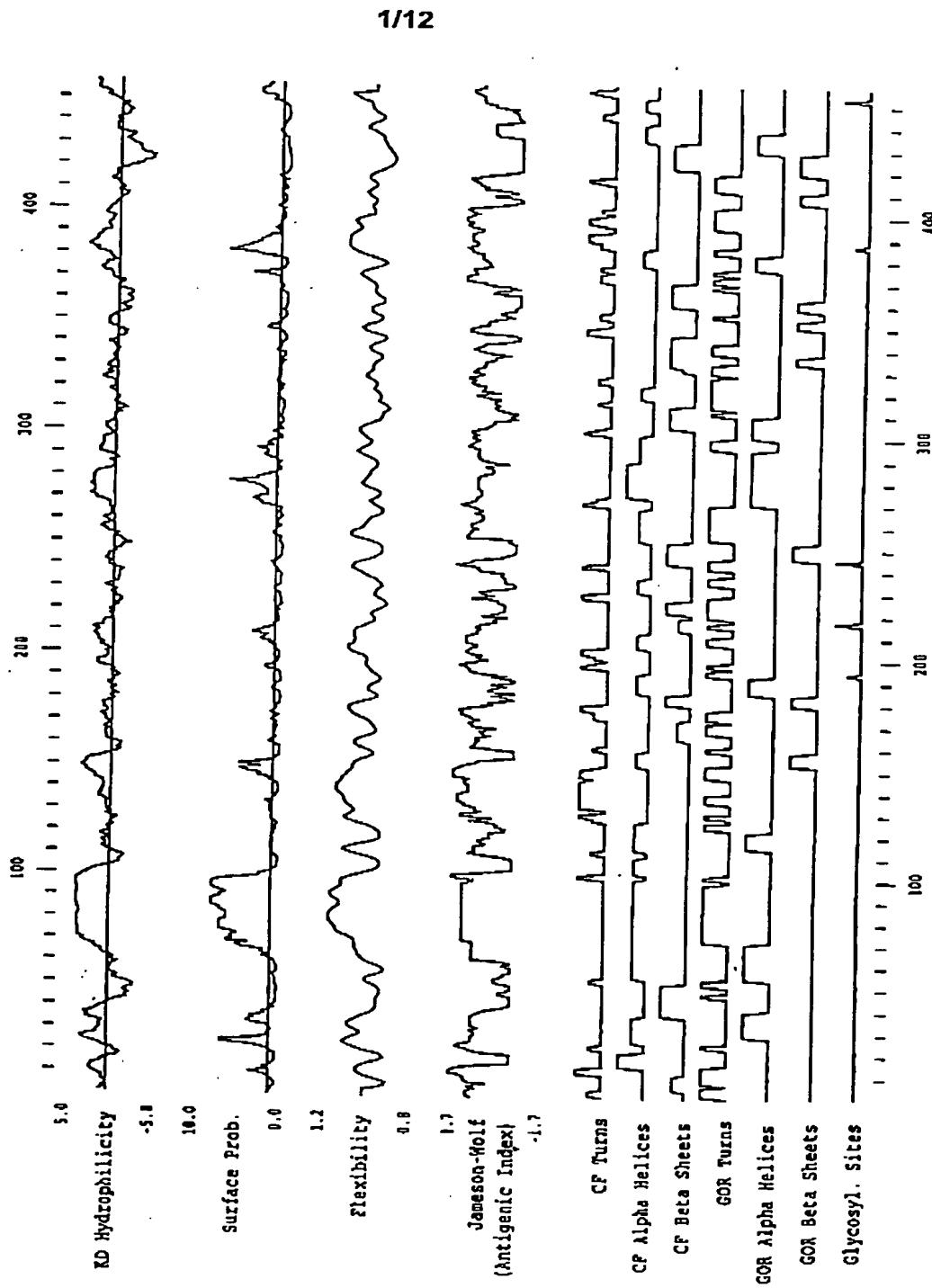
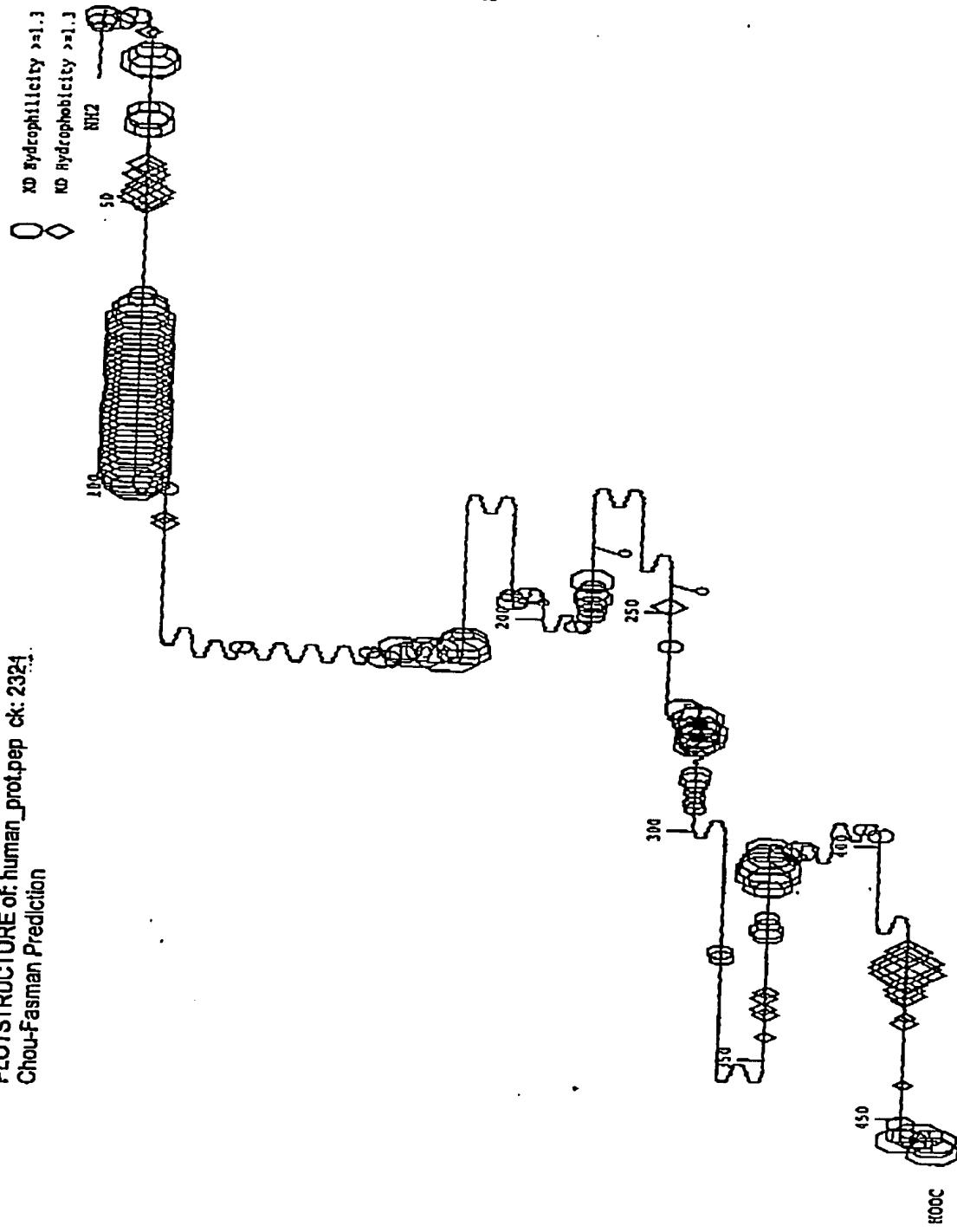


FIGURE 1

6 6 2 T C T " H B E S T H G D

PLOTSTRUCTURE of: human\_prot.pep ck: 2324  
Chou-Fasman Prediction



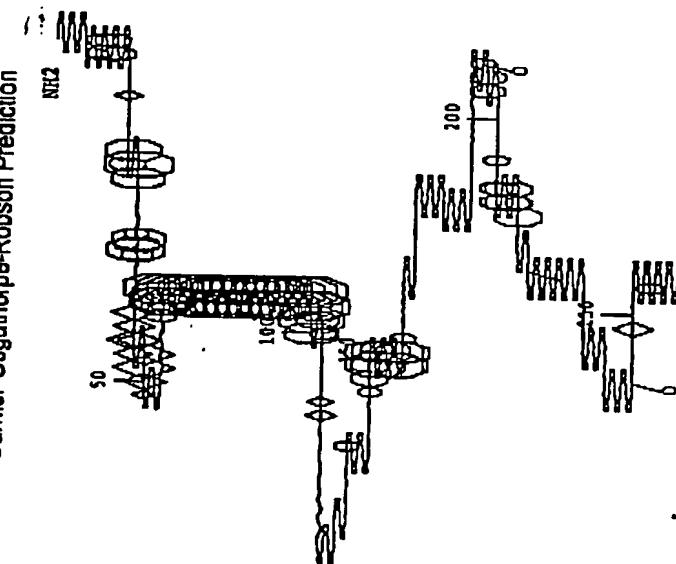
2/12

FIGURE 2

662707 " FIGURE 3 Thio

PLOTSTRUCTURE of: human\_prot.pep ck: 2321.  
Garnier-Osguthorpe-Robson Prediction

Hydrophilicity >1.3  
Hydrophobicity >1.3



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FIGURE 3

ESTIMATE PROTEIN

PEPTIDESTRUCTURE of: g713estmouse.0.12\_tr13.pop ck: 1880, 1 to 467

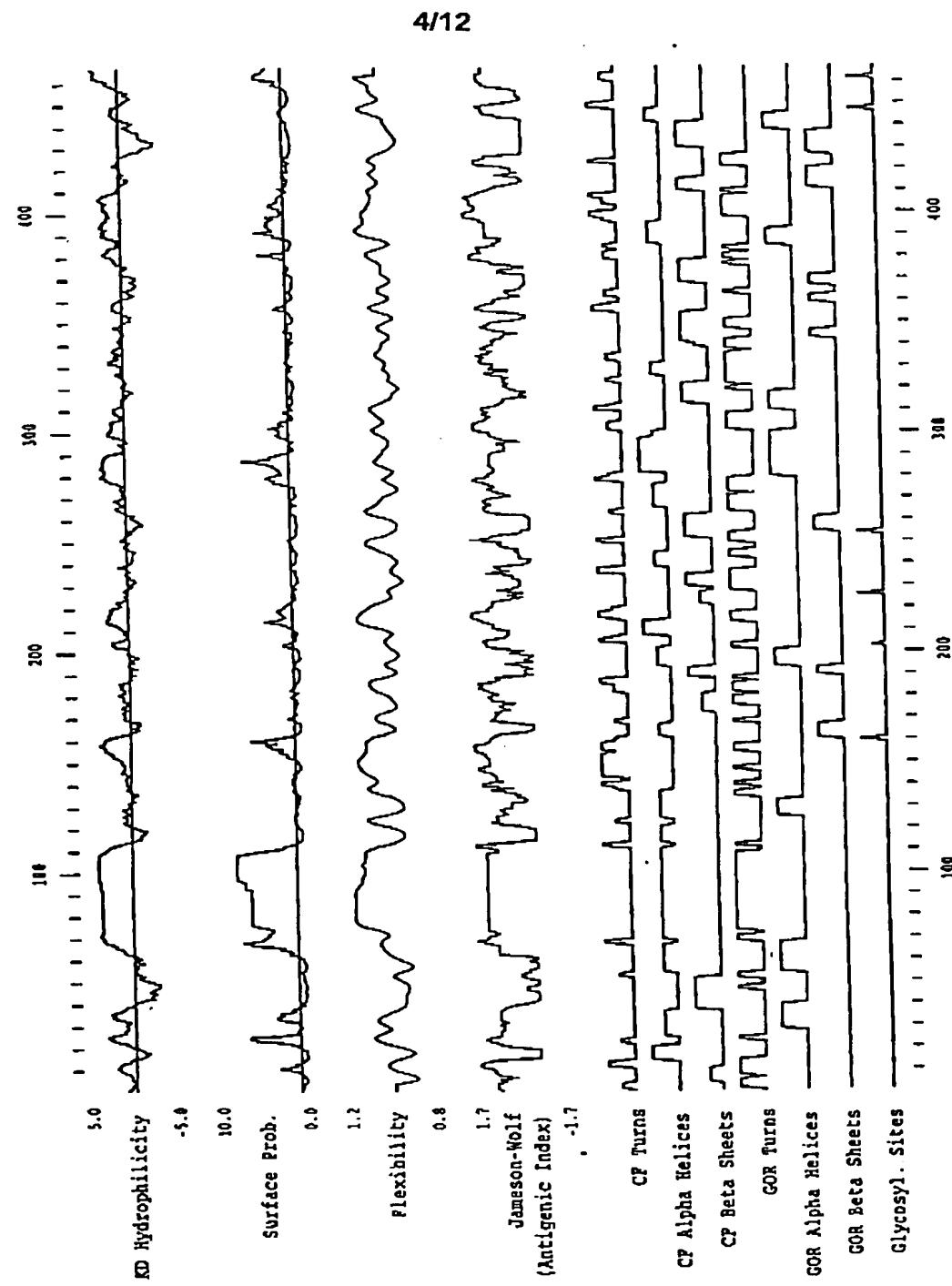


FIGURE 4

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PLOTSTRUCTURE of g713estmouse.0.12\_013:pep ck: 1880  
Chou-Fasman Prediction

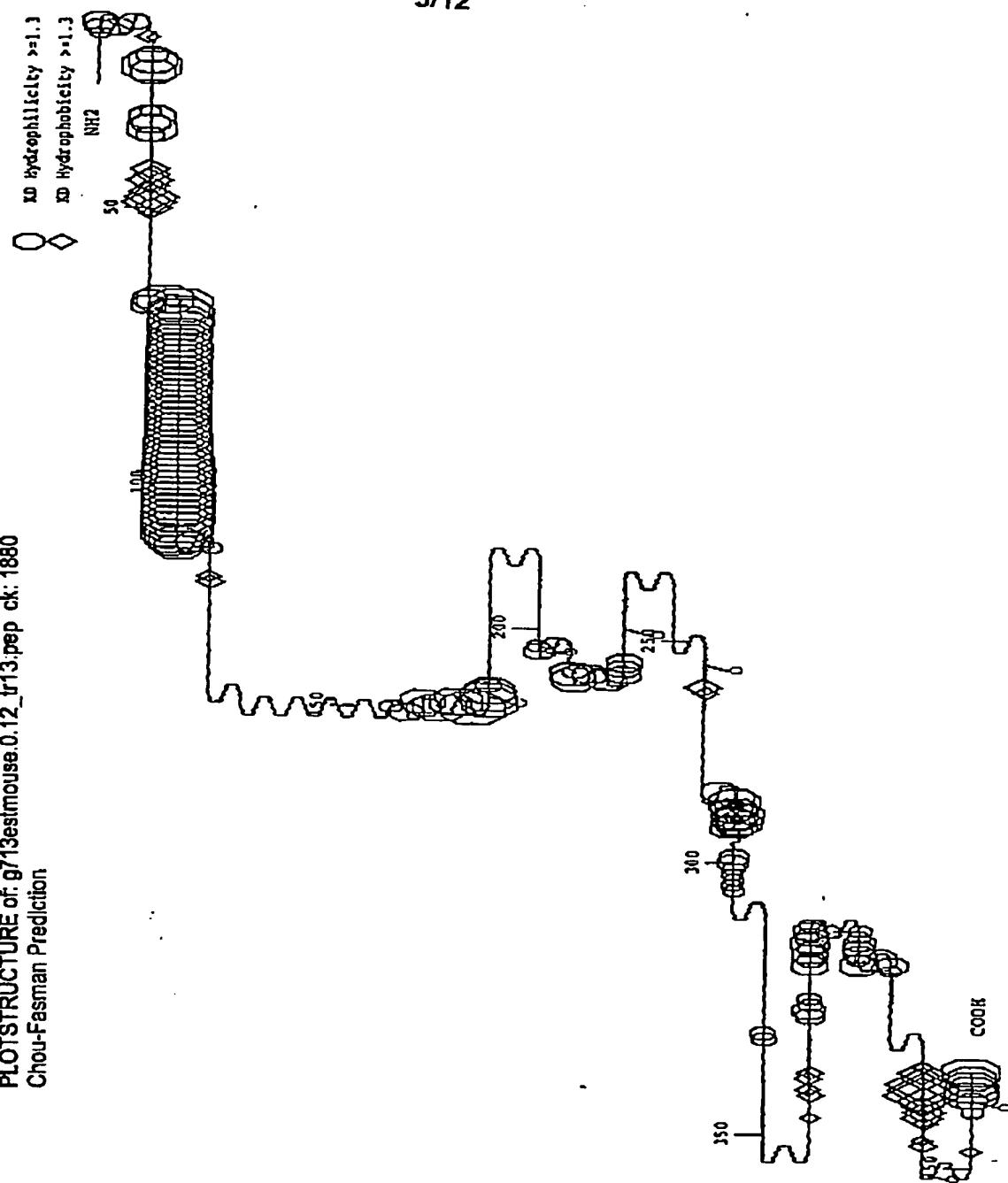
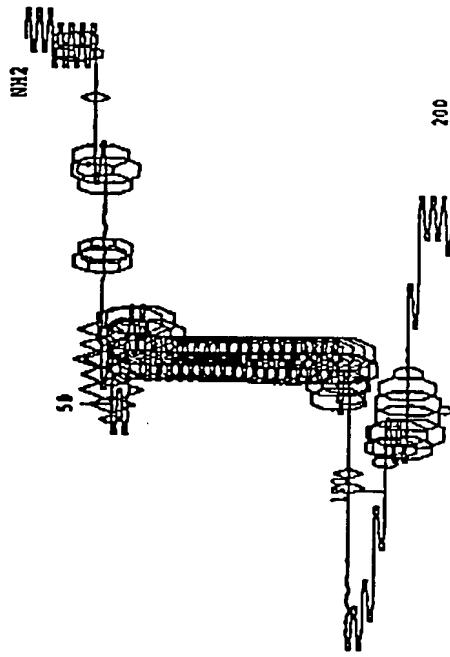


FIGURE 5

bioactive peptide

PLOTSTRUCTURE of: g713estmouse.0.12\_tr13.pep ck: 1880  
Garnier-Osguthorpe-Robson Prediction

Hydropathicity  $\alpha=1.1$   
Hydrophobicity  $\alpha=1.1$



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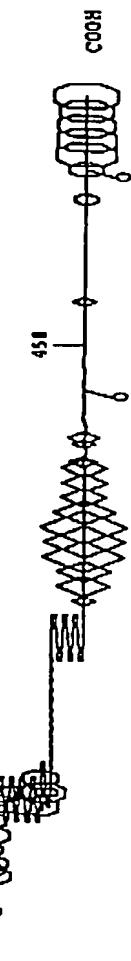
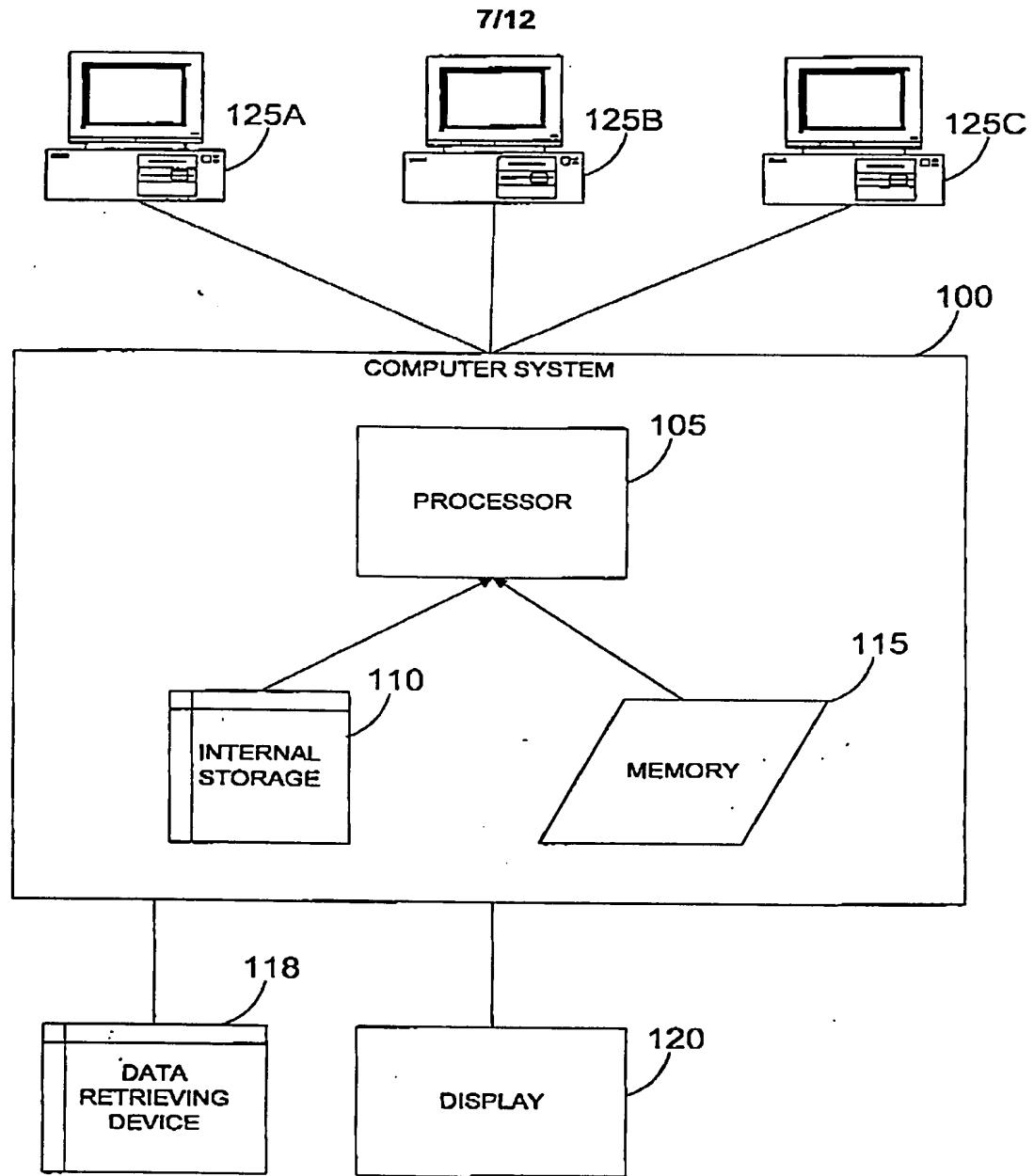


FIGURE 6

09416384-101299



**FIGURE 7**

00013284-2012090

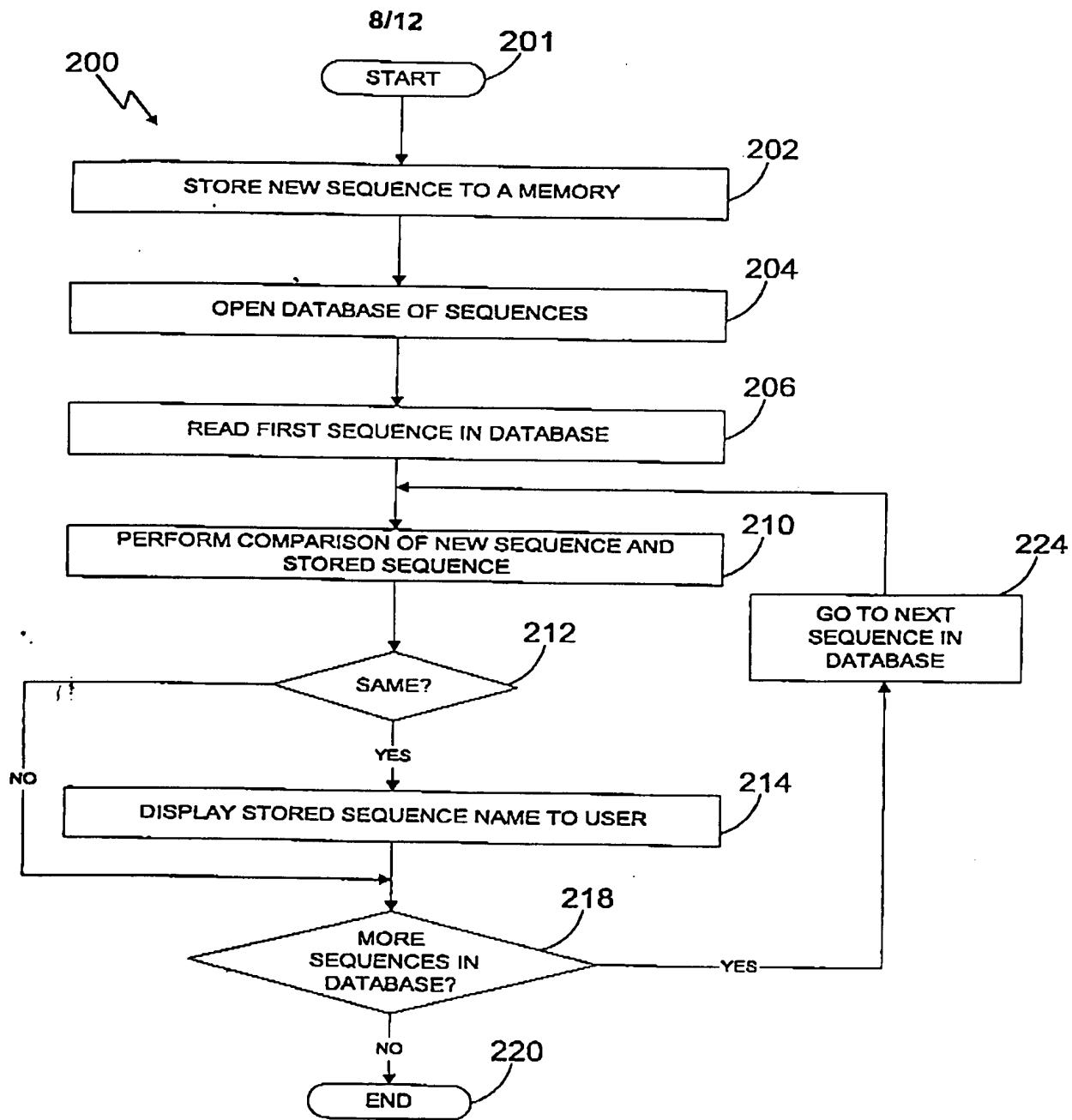


FIGURE 8

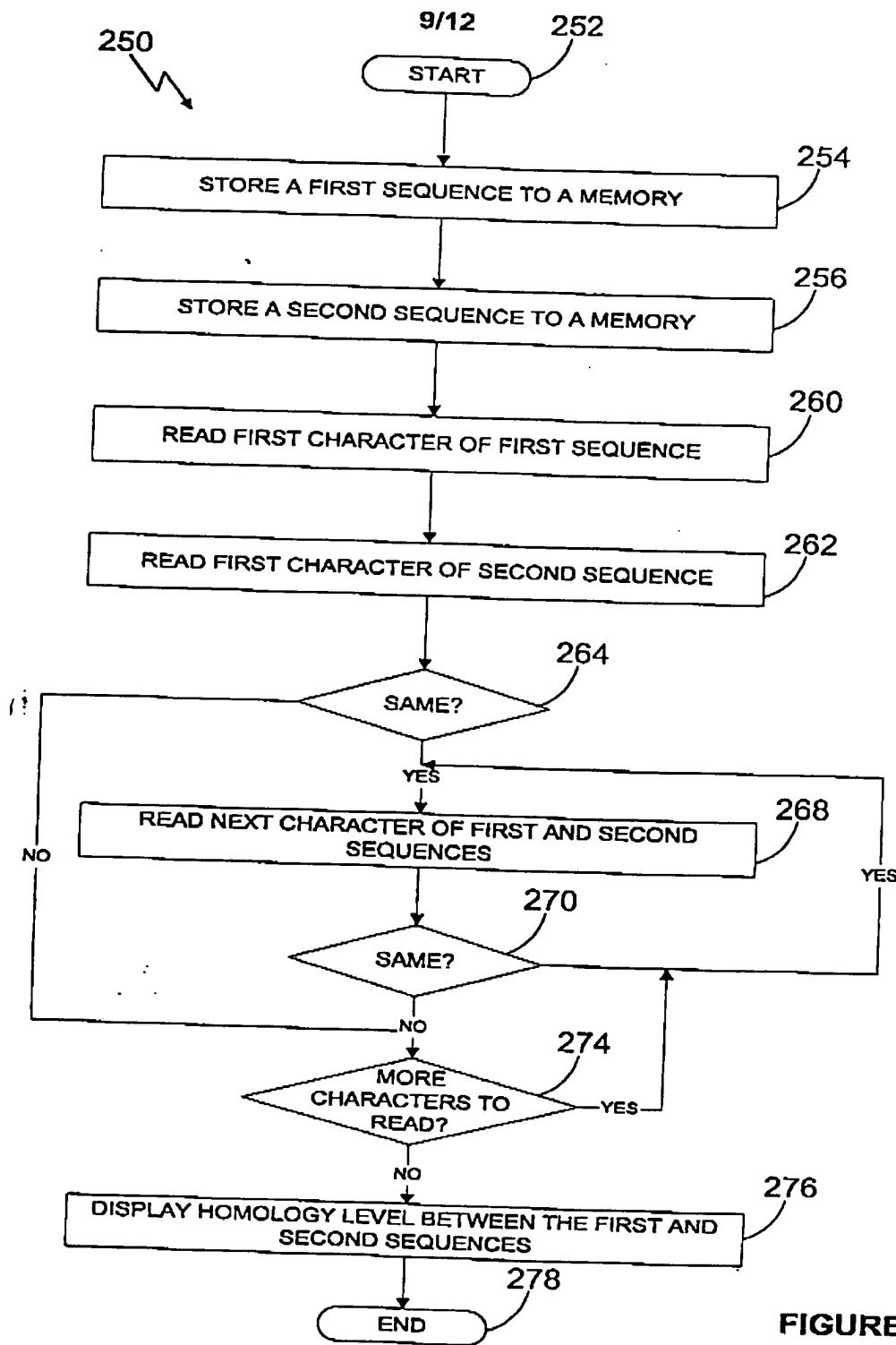


FIGURE 9

00453847-101200

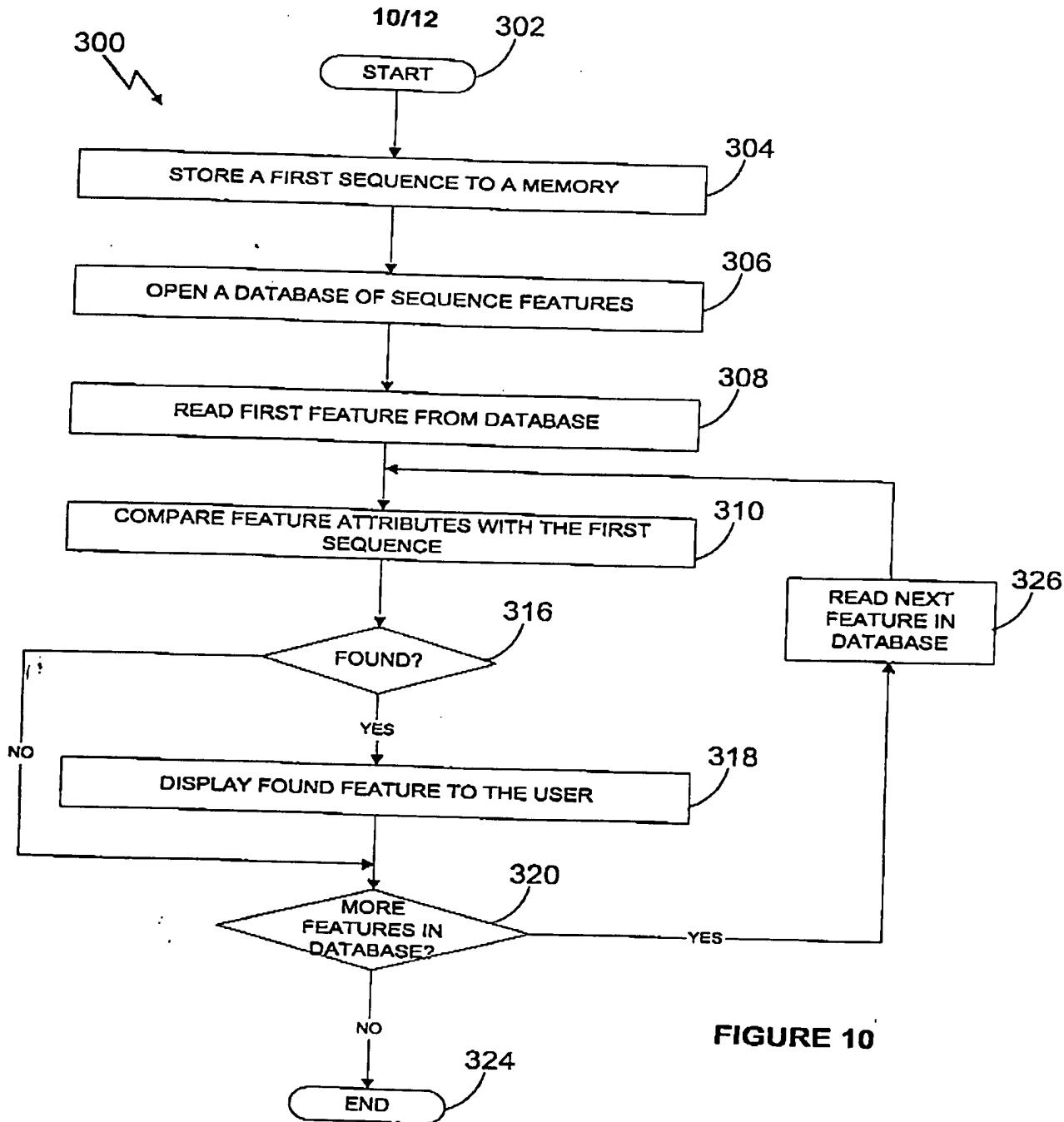


FIGURE 10

662701 "11825 Th1610

FIGURE 11

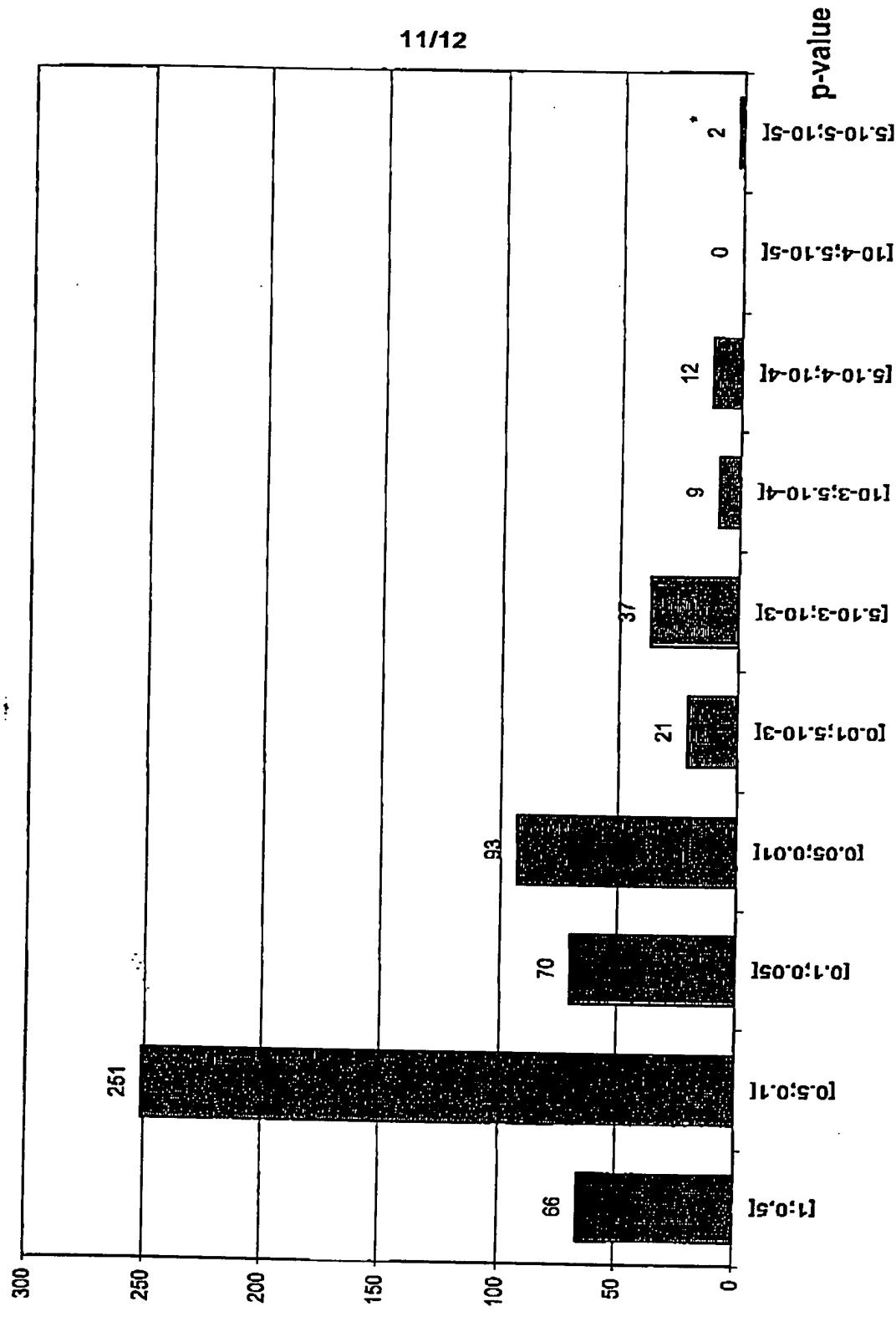


FIGURE 12

